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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/813,588

DATE: 09/02/2004
TIME: 07:43:28

Input Set : N:\Crf3\RULE60\10813588.raw.txt
Output Set: N:\CRF4\09022004\J813588.raw

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1 <110> APPLICANT: Donoho, Gregory
2     Turner, C. Alexander Jr.
3     Nehls, Michael
4     Friedrich, Glenn
5     Zambrowicz, Brian
6     Sands, Arthur T.
7 <120> TITLE OF INVENTION: Novel Human Proteins and Polynucleotides
8     Encoding the Same
9 <130> FILE REFERENCE: LEX-0071-USA
10 <140> CURRENT APPLICATION NUMBER: US/10/813,588
C--> 11 <141> CURRENT FILING DATE: 2004-03-30
12 <150> PRIOR APPLICATION NUMBER: US/09/691,344
13 <151> PRIOR FILING DATE: 2000-10-18
14 <150> PRIOR APPLICATION NUMBER: US 60/160,285
15 <151> PRIOR FILING DATE: 1999-10-19
16 <150> PRIOR APPLICATION NUMBER: US 60/183,583
17 <151> PRIOR FILING DATE: 2000-02-18
18 <160> NUMBER OF SEQ ID NOS: 7
19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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22 <211> LENGTH: 1464
23 <212> TYPE: DNA
24 <213> ORGANISM: homo sapiens
25 <400> SEQUENCE: 1
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28 acctgtgctt ctgactatct tctttcacc agctttcag atcaaatatgg tccatactgt 180
29 ggaagtatga ctgttccaa agaacttgg ttgaacacaa gtgaagtaac cgtccgctt 240
30 gagagtggat cccacatttc tggccgggtt ttttgctga cctatgcgag cagcgaccat 300
31 ccagatttaa taacatgttt ggaacgagct agccattatt tgaagacaga atacagcaa 360
32 ttctgccag ctgggttag agacgttagca ggagacattt ctggaaatat ggttagatgga 420
33 tatagagata cctcttattt gtcaaagct gccatccatg caggaataat tgctgtatgaa 480
34 ctaggtggcc agatcagtgt gttcagcgc aaaggatca gtcgatatga agggattctg 540
35 gccaatggtg ttcttcgag ggtatggttcc ctgtcagaca agcgatttct gtttacctcc 600
36 aatggttgca gcagatcctt gagtttgaa cctgacgggc aaatcagagc ttcttcctca 660
37 tggcagtcgg tcaatgagag tggagaccaa gttcactgggt ctcctggcca agccccactt 720
38 cagaccaag gcccacatcg ggcttcgggc gacagtagca acaaccacaa accacgagag 780
39 tggctggaga tcgatttggg ggagaaaaag aaaataacag gaattaggac cacaggatct 840
40 acacagtgcg acttcaactt ttatgttaag agtttgtga tgaacttcaa aaacaataat 900
41 tctaagtggc agacctataa agaattgtg aataatgaag aaaaggtgt tcagggtaac 960
42 tctaacttcc gggaccctgt gcaaaaacaat ttcatccctc ccatcgtggc cagatatgtg 1020
43 cgggttgc cccagacatg gcaccagagg atagccttga aggtggagct cattgggtgc 1080
44 cagattacac aaggtaatga ttcatgggtg tggcgcaga caagtcaaag caccagtgtt 1140

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ENTERED

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45 tcaactaaga aagaagatga gacaatcaca aggccatcc cctcgaaaga aacatccaca 1200
46 ggaataaaaca ttacaacggt ggctattcca ttgggtgcct ttgttgtcct ggtgtttgct 1260
47 ggaatgggga tccttcgcgc cttagaaag aagaagaaga aaggaagtcc gtatggatca 1320
48 gcggaggctc agaaaaacaga ctgttggaaag cagattaaat atcccttgc cagacatcag 1380
49 tcagctgagt ttaccatcag ctatgataat gagaaggaga tgacacaaaa gtttagatctc 1440
50 atcacaagtg atatggcagg ttaa 1464

52 <210> SEQ ID NO: 2
53 <211> LENGTH: 487
54 <212> TYPE: PRT
55 <213> ORGANISM: homo sapiens
56 <400> SEQUENCE: 2
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      Glu Lys Thr Ile Thr Val Pro Lys Gly Lys Arg Leu Ile Leu Arg Leu
      20          25           30
      Gly Asp Leu Asp Ile Glu Ser Gln Thr Cys Ala Ser Asp Tyr Leu Leu
      35          40           45
      Phe Thr Ser Ser Asp Gln Tyr Gly Pro Tyr Cys Gly Ser Met Thr
      50          55           60
      Val Pro Lys Glu Leu Leu Asn Thr Ser Glu Val Thr Val Arg Phe
      65          70           75          80
      Glu Ser Gly Ser His Ile Ser Gly Arg Gly Phe Leu Leu Thr Tyr Ala
      85          90           95
      Ser Ser Asp His Pro Asp Leu Ile Thr Cys Leu Glu Arg Ala Ser His
      100         105          110
      Tyr Leu Lys Thr Glu Tyr Ser Lys Phe Cys Pro Ala Gly Cys Arg Asp
      115         120          125
      Val Ala Gly Asp Ile Ser Gly Asn Met Val Asp Gly Tyr Arg Asp Thr
      130         135          140
      Ser Leu Leu Cys Lys Ala Ala Ile His Ala Gly Ile Ile Ala Asp Glu
      145         150          155          160
      Leu Gly Gly Gln Ile Ser Val Leu Gln Arg Lys Gly Ile Ser Arg Tyr
      165         170          175
      Glu Gly Ile Leu Ala Asn Gly Val Leu Ser Arg Asp Gly Ser Leu Ser
      180         185          190
      Asp Lys Arg Phe Leu Phe Thr Ser Asn Gly Cys Ser Arg Ser Leu Ser
      195         200          205
      Phe Glu Pro Asp Gly Gln Ile Arg Ala Ser Ser Ser Trp Gln Ser Val
      210         215          220
      Asn Glu Ser Gly Asp Gln Val His Trp Ser Pro Gly Gln Ala Arg Leu
      225         230          235          240
      Gln Asp Gln Gly Pro Ser Trp Ala Ser Gly Asp Ser Ser Asn Asn His
      245         250          255
      Lys Pro Arg Glu Trp Leu Glu Ile Asp Leu Gly Glu Lys Lys Lys Ile
      260         265          270
      Thr Gly Ile Arg Thr Thr Gly Ser Thr Gln Ser Asn Phe Asn Phe Tyr
      275         280          285
      Val Lys Ser Phe Val Met Asn Phe Lys Asn Asn Asn Ser Lys Trp Lys
      290         295          300

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95	Thr Tyr Lys Gly Ile Val Asn Asn Glu Glu Lys Val Phe Gln Gly Asn	
96	305	310 315 320
97	Ser Asn Phe Arg Asp Pro Val Gln Asn Asn Phe Ile Pro Pro Ile Val	
98	325	330 335
99	Ala Arg Tyr Val Arg Val Val Pro Gln Thr Trp His Gln Arg Ile Ala	
100	340	345 350
101	Leu Lys Val Glu Leu Ile Gly Cys Gln Ile Thr Gln Gly Asn Asp Ser	
102	355	360 365
103	Leu Val Trp Arg Lys Thr Ser Gln Ser Thr Ser Val Ser Thr Lys Lys	
104	370	375 380
105	Glu Asp Glu Thr Ile Thr Arg Pro Ile Pro Ser Glu Glu Thr Ser Thr	
106	385	390 395 400
107	Gly Ile Asn Ile Thr Thr Val Ala Ile Pro Leu Val Leu Leu Val Val	
108	405	410 415
109	Leu Val Phe Ala Gly Met Gly Ile Phe Ala Ala Phe Arg Lys Lys Lys	
110	420	425 430
111	Lys Lys Gly Ser Pro Tyr Gly Ser Ala Glu Ala Gln Lys Thr Asp Cys	
112	435	440 445
113	Trp Lys Gln Ile Lys Tyr Pro Phe Ala Arg His Gln Ser Ala Glu Phe	
114	450	455 460
115	Thr Ile Ser Tyr Asp Asn Glu Lys Glu Met Thr Gln Lys Leu Asp Leu	
116	465	470 475 480
117	Ile Thr Ser Asp Met Ala Gly	
118	485	
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121 <211>	LENGTH: 1761	
122 <212>	TYPE: DNA	
123 <213>	ORGANISM: homo sapiens	
124 <400>	SEQUENCE: 3	
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126	gaggcggccc ggccggggca gctgcggctc gggatccgtc gaggggaggg cgagttgcc	120
127	aagctggcgc ccagcggggt catggtgccc ggccggccgc gcggcgccgc actggcgcgg	180
128	gctgcggggc ggggcctcct ggcttgctg ctgcggctc ccgcggcgct ccggctgcag	240
129	gcggaggaggc tgggtgatgg ctgtggacac cttagtactt atcaggatag tggcacaatg	300
130	acatctaaga attatccgg gacctacccc aatcacactg tttgcgaaaa gacaattaca	360
131	gtaccaaagg ggaaaagact gattctgagg ttgggagatt tggatatcga atcccagacc	420
132	tgtgcttcgt actatcttct cttcaccaggc tcttcagatc aatatggtcc atactgtgga	480
133	agatgactg ttcccaaaga actcttggtt aacacaagtg aagtaacccgt ccgcggcgat	540
134	agtggatccc acatttctgg ccggggtttt ttgctgaccc atgcgagcag cgaccatcca	600
135	gatttaataa catgtttgga acgagctagc cattattga agacagaata cagcaaattc	660
136	tgcccagctg gttgttagaga cgtacggcggc gacatttctg ggaatatggt agatggat	720
137	agagataacct ctttattgtg caaagctgcc atccatgcag gaataattgc tgatgaacta	780
138	ggtggccaga tcagtgtgct tcagcgc当地 gggatcagtc gatatgaagg gattctggcc	840
139	aatgggttcc tttcggggca tggttccctg tcagacaaagg gatttctgtt tacctccat	900
140	ggttgcagca gatccttgc ttttgaacct gacggggcaaa tcagagcttc ttccatgg	960
141	cagtcggta atgagagtttgg agaccaagtt cactggctc ctggccaaac ccgacttcag	1020
142	gaccaaggcc catcatggcc ttccggccgac agtagcaaca accacaaacc acgagatgg	1080
143	ctggagatcg atttggggca gaaaaagaaa ataacaggaa ttaggaccac aggatctaca	1140
144	cagtcgaact tcaacttta tgttaagagt tttgtgatga acttcaaaaa caataattct	1200

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145	aagtggaga cctataaaagg aattgtgaat aatgaagaaa aggtgttca gggtaactct	1260
146	aaccttcggg acccagtgc aacaatttc atccctccca tcgtggccag atatgtgcgg	1320
147	gttgtcccc agacatggca ccagaggata gccttgaagg tggagctcat tggtgccag	1380
148	attacacaag gtaatgattc attgggtgtgg cgcaagacaa gtcaaagcac cagtgtttca	1440
149	actaagaaaag aagatgagac aatcacaagg cccatcccct cggaaagaaaac atccacagga	1500
150	ataaacatta caacgtggc tattccattt gtgctccttg ttgtcctgg tttgtctgga	1560
151	atggggatct ttgcagcctt tagaaagaag aagaagaaaag gaagtccgta tggatcagcg	1620
152	gaggctcaga aaacagactg ttggaagcag attaaatatc ccttgccag acatcagtca	1680
153	gctgagtttta ccatcagcta tgataatgag aaggagatga cacaaaagtt agatctcatc	1740
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159	<213> ORGANISM: homo sapiens	
160	<400> SEQUENCE: 4	
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163	Ser Ser Ala Glu Glu Ala Ala Arg Pro Gly Gln Leu Arg Leu Gly Ile	
164	20 25 30	
165	Arg Arg Gly Glu Ala Glu Leu Ala Lys Leu Ala Pro Ser Gly Val Met	
166	35 40 45	
167	Val Pro Gly Ala Arg Gly Gly Ala Leu Ala Arg Ala Ala Gly Arg	
168	50 55 60	
169	Gly Leu Leu Ala Leu Leu Ala Val Ser Ala Pro Leu Arg Leu Gln	
170	65 70 75 80	
171	Ala Glu Glu Leu Gly Asp Gly Cys Gly His Leu Val Thr Tyr Gln Asp	
172	85 90 95	
173	Ser Gly Thr Met Thr Ser Lys Asn Tyr Pro Gly Thr Tyr Pro Asn His	
174	100 105 110	
175	Thr Val Cys Glu Lys Thr Ile Thr Val Pro Lys Gly Lys Arg Leu Ile	
176	115 120 125	
177	Leu Arg Leu Gly Asp Leu Asp Ile Glu Ser Gln Thr Cys Ala Ser Asp	
178	130 135 140	
179	Tyr Leu Leu Phe Thr Ser Ser Asp Gln Tyr Gly Pro Tyr Cys Gly	
180	145 150 155 160	
181	Ser Met Thr Val Pro Lys Glu Leu Leu Leu Asn Thr Ser Glu Val Thr	
182	165 170 175	
183	Val Arg Phe Glu Ser Gly Ser His Ile Ser Gly Arg Gly Phe Leu Leu	
184	180 185 190	
185	Thr Tyr Ala Ser Ser Asp His Pro Asp Leu Ile Thr Cys Leu Glu Arg	
186	195 200 205	
187	Ala Ser His Tyr Leu Lys Thr Glu Tyr Ser Lys Phe Cys Pro Ala Gly	
188	210 215 220	
189	Cys Arg Asp Val Ala Gly Asp Ile Ser Gly Asn Met Val Asp Gly Tyr	
190	225 230 235 240	
191	Arg Asp Thr Ser Leu Leu Cys Lys Ala Ala Ile His Ala Gly Ile Ile	
192	245 250 255	
193	Ala Asp Glu Leu Gly Gly Gln Ile Ser Val Leu Gln Arg Lys Gly Ile	
194	260 265 270	

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195	Ser Arg Tyr Glu Gly Ile Leu Ala Asn Gly Val Leu Ser Arg Asp Gly			
196	275	280	285	
197	Ser Leu Ser Asp Lys Arg Phe Leu Phe Thr Ser Asn Gly Cys Ser Arg			
198	290	295	300	
199	Ser Leu Ser Phe Glu Pro Asp Gly Gln Ile Arg Ala Ser Ser Ser Trp			
200	305	310	315	320
201	Gln Ser Val Asn Glu Ser Gly Asp Gln Val His Trp Ser Pro Gly Gln			
202	325	330	335	
203	Ala Arg Leu Gln Asp Gln Gly Pro Ser Trp Ala Ser Gly Asp Ser Ser			
204	340	345	350	
205	Asn Asn His Lys Pro Arg Glu Trp Leu Glu Ile Asp Leu Gly Glu Lys			
206	355	360	365	
207	Lys Lys Ile Thr Gly Ile Arg Thr Thr Gly Ser Thr Gln Ser Asn Phe			
208	370	375	380	
209	Asn Phe Tyr Val Lys Ser Phe Val Met Asn Phe Lys Asn Asn Asn Ser			
210	385	390	395	400
211	Lys Trp Lys Thr Tyr Lys Gly Ile Val Asn Asn Glu Glu Lys Val Phe			
212	405	410	415	
213	Gln Gly Asn Ser Asn Phe Arg Asp Pro Val Gln Asn Asn Phe Ile Pro			
214	420	425	430	
215	Pro Ile Val Ala Arg Tyr Val Arg Val Val Pro Gln Thr Trp His Gln			
216	435	440	445	
217	Arg Ile Ala Leu Lys Val Glu Leu Ile Gly Cys Gln Ile Thr Gln Gly			
218	450	455	460	
219	Asn Asp Ser Leu Val Trp Arg Lys Thr Ser Gln Ser Thr Ser Val Ser			
220	465	470	475	480
221	Thr Lys Lys Glu Asp Glu Thr Ile Thr Arg Pro Ile Pro Ser Glu Glu			
222	485	490	495	
223	Thr Ser Thr Gly Ile Asn Ile Thr Thr Val Ala Ile Pro Leu Val Leu			
224	500	505	510	
225	Leu Val Val Leu Val Phe Ala Gly Met Gly Ile Phe Ala Ala Phe Arg			
226	515	520	525	
227	Lys Lys Lys Lys Gly Ser Pro Tyr Gly Ser Ala Glu Ala Gln Lys			
228	530	535	540	
229	Thr Asp Cys Trp Lys Gln Ile Lys Tyr Pro Phe Ala Arg His Gln Ser			
230	545	550	555	560
231	Ala Glu Phe Thr Ile Ser Tyr Asp Asn Glu Lys Glu Met Thr Gln Lys			
232	565	570	575	
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234	580	585		
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237	<211> LENGTH: 1620			
238	<212> TYPE: DNA			
239	<213> ORGANISM: homo sapiens			
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242	gctttgctgc tcgcggcttc cgccccggctc cggctgcagg cggaggagct gggtgatggc	120		
243	tgtggacacc tagtgactta tcaggatagt ggcacaatga catctaagaa ttatccggg	180		
244	acctacccca atcacactgt ttgcgaaaag acaattacag taccaaaggg gaaaagactg	240		

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/813,588

DATE: 09/02/2004

TIME: 07:43:29

Input Set : N:\Crf3\RULE60\10813588.raw.txt

Output Set: N:\CRF4\09022004\J813588.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date